

**IN THE SPECIFICATION**

*On page 11, please replace the paragraph from line 8 to line 11 with the following amended paragraph:*

In general, a preferred zinc finger framework has the structure:

(A)  $X_{0-2} C X_{1-5} C X_{9-14} H X_{3-6} H/C$  (SEQ ID NO:34)

where X is any amino acid, and the numbers in subscript indicated the possible numbers of residues represented by X.

*On pages 11 and 12, please replace the paragraph from line 12 on page 11 to line 2 on page 12 with the following amended paragraph:*

In a preferred aspect of the present invention, zinc finger nucleic acid binding motifs may be represented as motifs having the flowing primary structure:

(B)  $X^a C X_{2-4} C X_{2-3} F X^c X X X X L X X H X X X^b H$  - linker (SEQ ID NO: 22)

wherein X (including  $X^a$ ,  $X^b$  and  $X^c$ ) is any amino acid.  $X_{2-4}$  and  $X_{2-3}$  refer to the presence of 2 [[or]] to 4, or 2 or 3, amino acids, respectively, and  $X X X X L X X H X X$  between  $X^c$  and  $X^b$  are designated positions -1, 1, 2,3,4,5,6,7,8, and 9. The Cys and His residues, which together coordinate the zinc metal atom, are marked in bold text and are usually invariant, as is the Leu residue at position +4 in the  $\alpha$ -helix.